Fig. 1
(SEQ ID NO: 1)

```
-9    cggtgaagaatggtgtatttttaaatttcatgtcaataaccaatgtcccggtgctgaag
             M  V  Y  F  L  N  F  M  S  I  T  N  V  P  V  L  K
52    cgcgcgcgactctacatggcgacgaatcgccggctggtggttgttcttgtggtgctgctg
       R  A  R  L  Y  M  A  T  N  R  R  L  V  V  V  L  V  V  L  L
112   tactgggtggtccagaacgtttggacgtggacgcctgggacgcgcgatttggcccaagtg
       Y  W  V  V  Q  N  V  W  T  W  S  P  G  T  R  D  L  A  Q  V
172   gacgcgaagatcgaggccgagctaaactcgaatctacatacttttggagcgcatttcgcgc
       D  A  K  I  E  A  E  L  N  S  N  L  H  T  F  G  A  H  L  R
232   cacttaaaccggcttccggcagagtcggccaccctgcgtgaaaaactcaccttctatttc
       H  L  N  R  L  P  A  E  S  A  T  L  R  E  K  L  T  F  Y  F
282   ccatattatcctgaaaagcccgtgccgaaccagatctggcagacatggaaggtcgatctc
       P  Y  Y  P  E  K  P  V  P  N  Q  I  W  Q  T  W  K  V  D  L
352   gaagacgacaacttccccaagcagtacagacggtttcagaagacgtgggtcgagaaaaat
       E  D  D  N  F  P  K  Q  Y  R  R  F  Q  K  T  W  V  E  K  N
412   ccagactacgtgtaccacctgattccggactctgtgattgaggactttgtggcgagtttg
       P  D  Y  V  Y  H  L  I  P  D  S  V  I  E  D  F  V  A  S  L
472   tacgcgaacgtgccggaggtggtcagagcgtaccagctgcttccgaaaaatatcatgaag
       Y  A  N  V  P  E  V  V  R  A  Y  Q  L  L  P  K  N  I  M  K
532   gcggattttttccggtatttggtgatctacgcgcgcggaggcacctactcagacatggac
       A  D  F  F  R  Y  L  V  I  Y  A  R  G  G  T  Y  S  D  M  D
592   acggtgtgtttaaagccgatcaaggactgggccacgtttgatcgcgacctgatccacgct
       T  V  C  L  K  P  I  K  D  W  A  T  F  D  R  D  L  I  H  A
652   gccgacaataaggccgatctctcccagatagatccggaagcaagaaccacgcctgtgggg
       A  D  N  K  A  D  L  S  Q  I  D  P  E  A  R  T  T  P  V  G
712   ctggtgattggcattgaggccgacccggacaggccgactggcacgagtggttctcgcgc
       L  V  I  G  I  E  A  D  P  D  R  P  D  W  H  E  W  F  S  R
772   agactgcagttctgccagtggacgatccaggcgaagccgggacaccgctgctgcgcgag
       R  L  Q  F  C  Q  W  T  I  Q  A  K  P  G  H  P  L  L  R  E
832   ctgatcatccggatcgtggaggagacgttccgcaaacagcacatgggcgttttgaaaaga
       L  I  I  R  I  V  E  E  T  F  R  K  Q  H  M  G  V  L  K  R
892   gtggaaggcaaggactcgggcgcagatatcatgcagtggacaggacgcgggatatttaca
       V  E  G  K  D  S  G  A  D  I  M  Q  W  T  G  P  G  I  F  T
952   gacactctgtttgattatctgaacaatgtggcgagcgacggcaagtgggcgacgggtac
       D  T  L  F  D  Y  L  N  N  V  A  S  D  G  K  L  G  D  G  Y
1012  ggcgtggggtcgttgtattggcgcaagcacggcaaatataagctgaaaaagacagaaatt
       G  V  G  S  L  Y  W  R  K  H  G  K  Y  K  L  K  K  T  E  I
1072  aacaagaataacgagccattgcattcgaggaccagcttatcaactggaggtcgctgacc
       N  K  N  N  E  P  L  H  S  E  D  Q  L  I  N  W  R  S  L  T
1132  aacatggacaagccaaagatcatgggggacgtaatggtgttaccaatcacgagctttagt
       N  M  D  K  P  K  I  M  G  D  V  M  V  L  P  I  T  S  F  S
1192  ccgaacgtggggcacatgggctcaaagagcagctcgataggctggcatttgtggagcat
       P  N  V  G  H  M  G  S  K  S  S  S  D  R  L  A  F  V  E  H
1252  ttattttctggcagctggaagccaaaaaacaaataggaaaaataaataattagctgcatt
       L  F  S  G  S  W  K  P  K  N  K  -
1312  ttagataattctcatgagcaggcacagaacg
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Fig. 2

SEQ ID NO.: